Amendments to the Claims

- 1. (previously presented) A *B. oleracea* plant resistant to clubroot disease, wherein the resistance is obtained from a clubroot resistant *B. rapa* plant and the resistance to clubroot is monogenic and dominant.
- 2. (original) The plant according to claim 1, wherein said *B. oleracea* plant is rated at level 2 or less in a test for the disease having a 1-9 scale or at a level 1 or less in a test for the disease having a 0-5 scale.
- 3. (original) The plant according to claim 1, wherein said *B. oleracea* plant is rated at level 1 in a test for the disease having a 1-9 scale or at a level 0 in a test for the disease having a 0-5 scale.
- 4. (original) The plant according to claim 1, wherein said *B. oleracea* plant is broccoli, white cabbage, cauliflower, Brussels sprouts, Borecole, Savoy, or red cabbage.
- 5. (original) The plant according to claim 1, wherein said resistance is linked to a molecular marker obtainable by PCR amplification.
- 6. (original) The plant according to claim 1, wherein said resistance is linked to a molecular marker obtainable by PCR amplification using primer O20 (SEQ ID NO:1) or primer Y13 (SEQ ID NO:2).
- 7. (original) The plant according to claim 5, wherein said resistance is within 10 cM of said molecular marker.
- 8. (original) The plant according to claim 5, wherein said resistance is within 6 cM of said molecular marker.
- 9. (cancelled)
- 10. (original) The plant according to claim 1, wherein said resistance is obtainable from Chinese cabbage F1 hybrid Parkin.
- 11. (previously presented) A *B. oleracea* plant comprising a locus conferring resistance to clubroot disease, wherein said resistance is obtained from a clubroot resistant *B. rapa* plant and said resistance is monogenic and dominant.

Claims 12-16. (cancelled)

- 17. (previously presented) The plant according to claim 1, wherein said *B. oleracea* plant is homozygous for said resistance.
- 18. (previously presented) The plant according to claim 1, wherein said *B. oleracea* plant is heterozygous for said resistance.
- 19. (previously presented) The plant according to claim 1, wherein said *B. oleracea* plant is an inbred or a dihaploid.
- 20. (previously presented) The plant according to claim 1, wherein said *B. oleracea* plant is a hybrid.
- 21. (previously presented) The plant according to claim 19, wherein said *B. oleracea* plant is cytoplasmic male sterile.
- 22. (previously presented) A seed of the plant according to claim 1.
- 23. (previously presented) A fruit or a part of the plant according to claim 1.
- 24. (previously presented) A part of the plant according to claim 1, wherein said part is pollen, ovule or embryo.

Claims 25-26. (cancelled)

- 27. (withdrawn) A method for producing a *B. oleracea* plant comprising a monogenic and dominant resistance to clubroot comprising the steps of:
 - a) obtaining a B. rapa plant resistant to clubroot:
 - b) crossing said B. rapa plant with a B. oleracea plant,
 - c) rescuing embryos resulting from the cross of step b):
 - d) regenerating a plant from a embryo of step c);
 - e) selecting a plant of step d) that is resistant to clubroot;
 - f) back-crossing a plant resulting from step e) with a B. oleracea plant.

- 28. (withdrawn) The method according to claim 27, further comprising introgressing the resistance into an elite *B. oleracea* inbred.
- 29. (withdrawn) The method according to claim 28, further comprising crossing said inbred to another *B. oleracea* inbred to produce a hybrid.
- 30. (withdrawn) A B. oleracea plant obtainable by the method of claim 27.
- 31. (withdrawn) A method for transferring a monogenic and dominant resistance to clubroot to a *B. oleracea* plant susceptible or less resistant to the disease comprising the steps of:
 - a) obtaining a *B. oleracea* plant comprising a monogenic and dominant resistance to clubroot;
 - b) crossing said *B. oleracea* plant of step a) with a *B. oleracea* plant susceptible or less resistant to clubroot;
 - c) selecting a plant from the cross of step b) that is resistant to clubroot.
- 32. (withdrawn) The method according to claim 31, further comprising backcrossing said resistance into said *B. oleracea* plant susceptible or less resistant to clubroot.
- 33. (withdrawn) A DNA fragment amplified from a Brassica genome, wherein said DNA fragment is approximately 400 bp long and comprises SEQ ID NO:1 or wherein said DNA fragment is approximately 640 bp long and comprises SEQ ID NO:2.
- 34. (cancelled)
- 35. (withdrawn) The DNA fragment according to claim 33, wherein said DNA fragment is indicative of the presence of a dominant and monogenic resistance to clubroot in a Brassica plant.

Claims 36-40. (cancelled)

41. (withdrawn) A kit for detecting a monogenic and dominant resistance to clubroot in a *B. oleracea* plant comprising an oligonucleotide set forth in SEQ ID NO:1 or SEQ ID NO:2.

- 42. (withdrawn) A method for transferring a monogenic and dominant resistance to clubroot to a *B. oleracea* plant susceptible or less resistant to the disease comprising the steps of:
 - a) obtaining a *B. oleracea* plant comprising a monogenic and dominant resistance to clubroot;
 - b) crossing said *B.oleracea* plant of step a) with a *B. oleracea* plant susceptible or less resistant to clubroot;
 - c) selecting a plant comprising a DNA fragment obtainable by PCR amplification using primer O20 (SEQ ID NO:1) or primer Y13 (SEQ ID NO:2);

wherein said plant of step c) is resistant to clubroot.

43. (withdrawn) The method according to claim 42, further comprising backcrossing said resistance into said *B. oleracea* plant susceptible or less resistant to clubroot.

Claims 44-45. (cancelled)

- 46. (previously presented) A seed of the plant according to claim 11.
- 47. (previously presented) A fruit or a part of the plant according to claim 11.
- 48. (previously presented) The plant according to claim 11, further comprising a nucleic acid sequence, which can be amplified by PCR using primer O20 (SEQ ID NO:1) or primer Y13 (SEQ ID NO:2).
- 49. (previously presented) The plant according to claim 48, wherein a DNA fragment of approximately 400 bp is amplified using primer O20 (SEQ ID NO:1).
- 50. (withdrawn) The plant according to claim 48, wherein a DNA fragment of approximately 640 bp is amplified using primer Y13 (SEQ ID NO:2).
- 51. (previously presented) A seed of the plant according to claim 48.
- 52. (previously presented) A fruit or a part of the plant according to claim 48.

- 53. (new) A *B. oleracea* plant resistant to clubroot disease, wherein the resistance is obtained from a clubroot resistant *B. rapa* plant, wherein the clubroot resistant *B. rapa* plant is a Chinese cabbage F1 hybrid 'Parkin' and the resistance to clubroot is monogenic and dominant.
- 54. (new) A *B. oleracea* plant resistant to clubroot disease wherein the resistance to clubroot is monogenic and dominant and wherein the plant is a plant of line CFL667 deposited with NCIMB under accession number NCIMB 41134, or a progeny or ancestor of said line CFL667 comprising the monogenic and dominant resistance to clubroot comprised in said line CFL667, or a plant derived from said line CFL667 deposited with NCIMB under accession number NCIMB 41134 and comprising the monogenic and dominant resistance to clubroot comprised in said line CFL667.